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CCCATTGAGGCTGTTGATCCCGAAGTGAAGACTGAGGTCTTCGCTGACTCCCTCCTTGCTGCAGCAGGCGATGACGACTGGGAGTCACCT  180
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AGCCCTGGTCCTACTTTCAATGTTCCCAGAGGAACAGAGACTGTAGTTAGGTTCATCAACAATGCCACCGTGGAGAACTCGGTCCATCTG  450
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CACGGCTCCCCATCGCGTGCCCCTTTCGATGGTTGGGCTGAAGATGTGACCTTCCCTGGGGAGTACAAGGATTACTACTTTCCCAACTAC  540
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CAATCCGCCCGCCTTCTGTGGTACCATGACCACGCTTTCATGAAGACTGCTGAGAATGCCTACTTTGGTCAGGCTGGCGCCTACATTATC  630
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GATGGTACCCTGCGTTCGACCGAGGGTGAGGACCAGGACCTGTGGGGAGATGTCATCCATGTCAACGGACAGCCATGGCCTTTCCTTAAC  810
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GCCCGAGTGCAGGAGCTGGCCGAGCAGGAGCCGTACAACCGCCTCGATGAGATCCTGGAGGATCTTGGAATCGAGGAGTAA          1791
 A  R  V  Q  E  L  A  E  Q  E  P  Y  N  R  L  D  E  I  L  E  D  L  G  I  E  E              594
```

Figure 1

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Figure 2

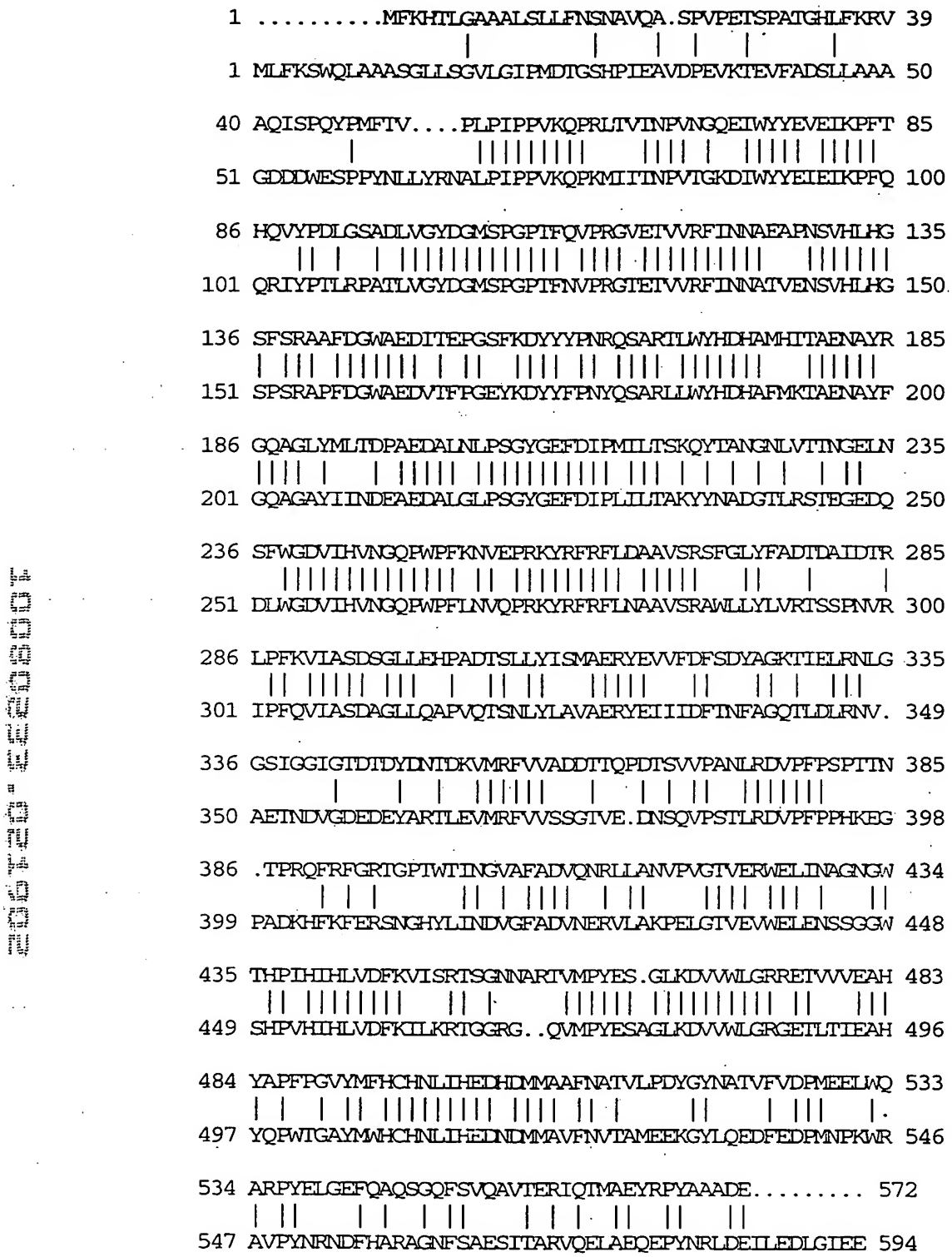
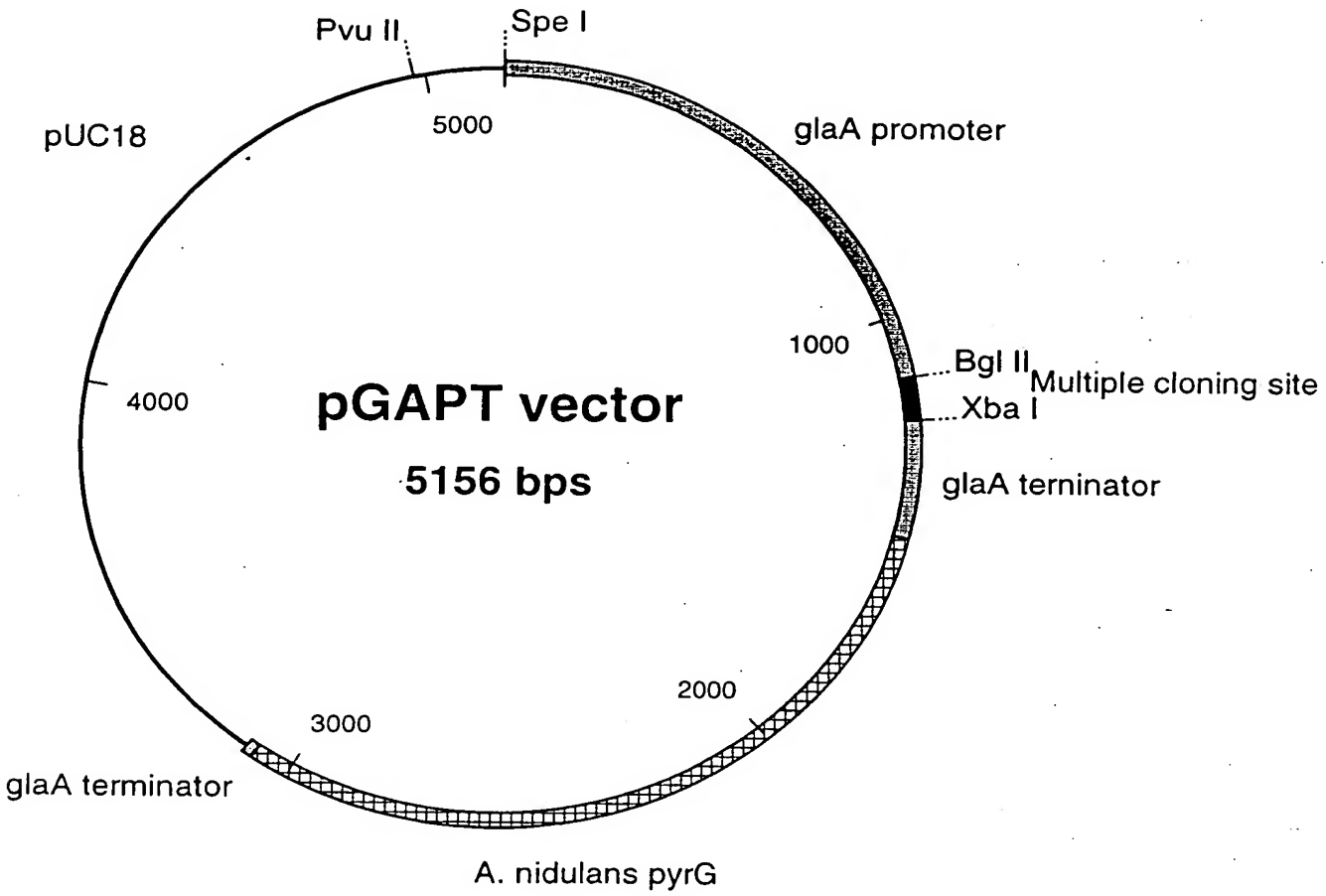


Figure 3: protein sequneces aligment of Bilirubin oxidase (top sequence) with Stachybotrys oxidase (bottom sequence).

Figure 3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2000
3000
glaA terminator
A. nidulans pyrG

Figure 4

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GTCCTCGGCA TCCCGATGGA CACCGGCAGC CACCCCATTG AGGCTGTTGA TCCCGAAGTG 120
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CCTCCATACA ACTTGCTTTA CAGGTGAGAC ACCTGTCCCA CCTGTTTTCC CTCGATAACT 240
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Figure 5

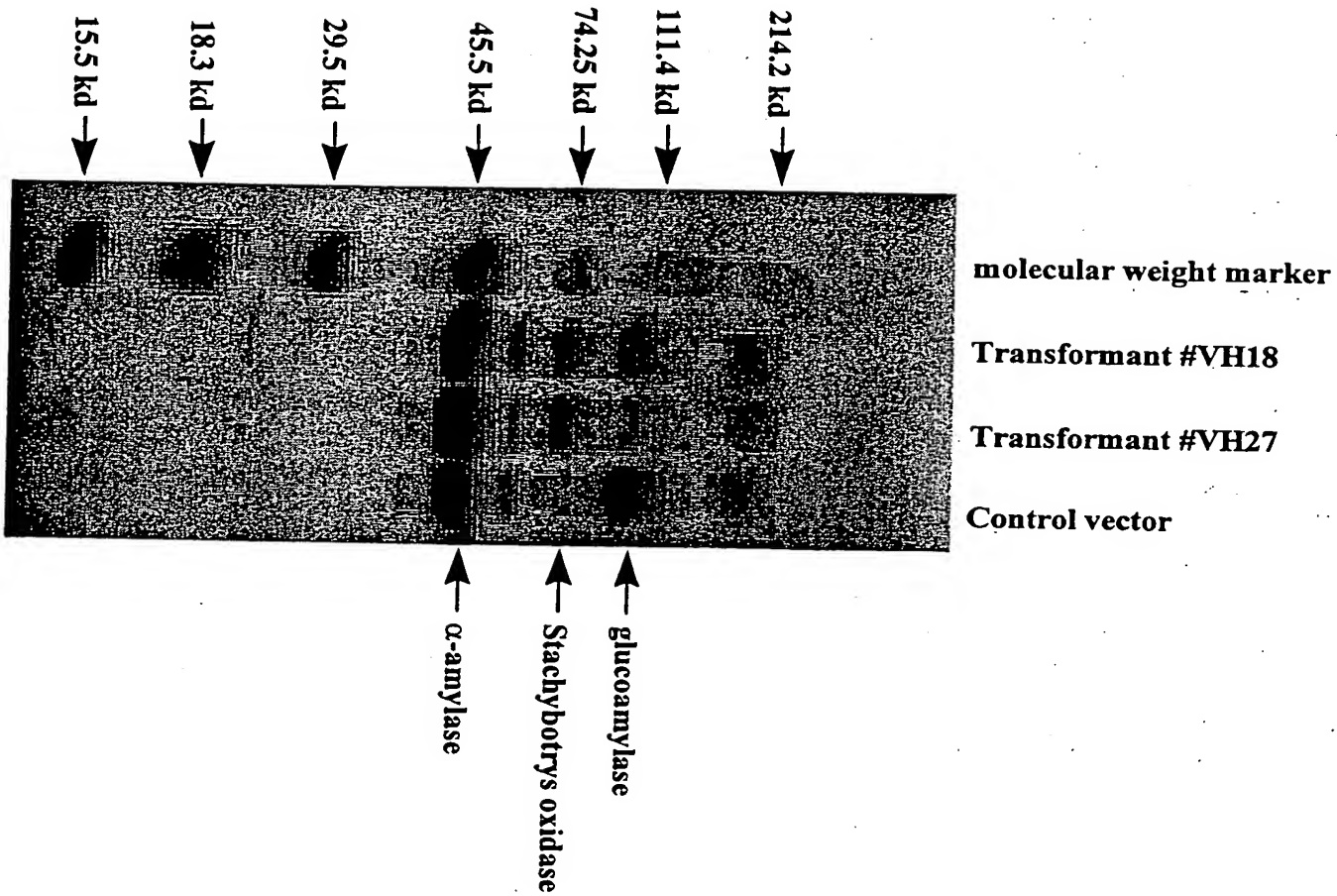
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Figure 6